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RAW SEQUENCE LISTING DATE: 07/18/2002 PATENT APPLICATION: US/09/931,009A TIME: 10:30:18

Input Set : N:\Crf3\07092002\I931009.raw
Output Set: N:\CRF3\07182002\I931009A.raw

1 <110> APPLICANT: Smith, Theresa H.

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2 <120> TITLE OF INVENTION: PRO-INFLAMMATORY FIBRINOPEPTIDE
 3 <130> FILE REFERENCE: US 1257/01 (VA)
 4 <140> CURRENT APPLICATION NUMBER: US/09/931,009A
 5 <141> CURRENT FILING DATE: 2002-06-28
 6 <160> NUMBER OF SEQ ID NOS: 2
 8 <210> SEQ ID NO: 1
 9 <211> LENGTH: 620
10 <212> TYPE: PRT
11 <213> ORGANISM: Homo sapiens
12 <220> FEATURE:
13 <400> SEQUENCE: 1
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14
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15
         Arg Gly Pro Arg Val Val Glu Arg His Gln Ser Ala Cys Lys Asp
16
17
         Ser Asp Trp Pro Phe Cys Ser Asp Glu Asp Trp Asn Tyr Lys Cys
18
19
         Pro Ser Gly Cys Arg Met Lys Gly Leu Ile Asp Glu Val Asn Gln
20
21
22
         Asp Phe Thr Asn Arg Ile Asn Lys Leu Lys Asn Ser Leu Phe Glu
                                                                    75
23
                          65
                                               70
         Tyr Gln Lys Asn Asn Lys Asp Ser His Ser Leu Thr Thr Asn Ile
24
25
                                               85
         Met Glu Ile Leu Arg Gly Asp Phe Ser Ser Ala Asn Asn Arg Asp
26
                                              100
27
                          95
         Asn Thr Tyr Asn Arg Val Ser Glu Asp Leu Arg Ser Arg Ile Glu
28
                                              115
29
                          110
         Val Leu Lys Arg Lys Val Ile Glu Lys Val Gln His Ile Gln Leu
30
                                              130
                         125
31
         Leu Gln Lys Asn Val Arg Ala Gln Leu Val Asp Met Lys Arg Leu
32
                                                                   150
33
                         140
                                              145
         Glu Val Asp Ile Asp Ile Lys Ile Arg Ser Cys Arg Gly Ser Cys
34
                          155
                                              160
35
         Ser Arg Ala Leu Ala Arg Glu Val Asp Leu Lys Asp Tyr Glu Asp
36
37
                                              175
         Gln Gln Lys Gln Leu Glu Gln Val Ile Ala Lys Asp Leu Leu Pro
38
39
                                              190
                                                                   195
                         185
         Ser Arg Asp Arg Gln His Leu Pro Leu Ile Lys Met Lys Pro Val
40
                                              205
41
                          200
         Pro Asp Leu Val Pro Gly Asn Phe Lys Ser Gln Leu Gln Lys Val
42
43
                         215
                                              220
         Pro Pro Glu Trp Lys Ala Leu Thr Asp Met Pro Gln Met Arg Met
44
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45					230					235					240
46	Glu	Leu	Glu	Arg	Pro	Gly	Gly	Asn	Glu	Ile	Thr	Arg	Gly	Gly	Ser
47					245					250					255
48	Thr	Ser	Tyr	Gly	Thr	Gly	Ser	Glu	Thr	Glu	Ser	Pro	Arg	Asn	Pro
49			_		260					265					270
50	Ser	Ser	Ala	Gly	Ser	Trp	Asn	Ser	Gly	Ser	Ser	Gly	${\tt Pro}$	Gly	Ser
51				_	275					280					285
52	Thr	Gly	Asn	Arg	Asn	Pro	Gly	Ser	Ser	Gly	Thr	Gly	Gly	Thr	Ala
53		_			290					295					300
54	Thr	Trp	Lys	Pro	Gly	Ser	Ser	Gly	Pro	Gly	Ser	Thr	Gly	Ser	${\tt Trp}$
55		_	_		305	,				310					315
56	Asn	Ser	Gly	Ser	Ser	Gly	Thr	Gly	Ser	Thr	Gly	Asn	Gln	Asn	Pro
57			-		320					325					330
58	Gly	Ser	Pro	Arg	Pro	Gly	Ser	Thr	Gly	Thr	Trp	Asn	${\tt Pro}$	Gly	Ser
59	1			_	335	-				340					345
60	Ser	Glu	Arq	Gly	Ser	Ala	Gly	His	${\tt Trp}$	Thr	Ser	Glu	Ser	Ser	Val
61			_	•	350		-		_	355					360
62	Ser	Glv	Ser	Thr	Gly	Gln	Trp	His	Ser	Glu	Ser	Gly	Ser	Phe	Arg
63					365		-			370					375
64	Pro	Asp	Ser	Pro	Gly	Ser	Gly	Asn	Ala	Arg	Pro	Asn	Asn	Pro	Asp
65					380		-			385					390
66	Trp	Glv	Thr	Phe	Glu	Glu	Val	Ser	Gly	Asn	Val	Ser	Pro	Gly	Thr
67		1			395				-	400					405
68	Ara	Ara	Glu	Tvr	His	Thr	Glu	Lys	Leu	Val	Thr	Ser	Lys	Gly	Asp
69	5	5	-	- 4	410			-		415				-	420
70	Lvs	Glu	Leu	Ara	Thr	Glv	Lys	Glu	Lys	Val	Thr	Ser	Gly	Ser	Thr
71					425	. 1	-		-	430					435
72	Thr	Thr	Thr	Arq	Arq	Ser	Cys	Ser	Lys	Thr	Val	Thr	Lys	Thr	Val
73				,	440		-		-	445					450
74	Ile	Gly	Pro	Asp	Gly	His	Lys	Glu	Val	Thr	Lys	Glu	Val	Val	Thr
75		. 1		-	455		-			460					465
76	Ser	Glu	Asp	Gly	Ser	Asp	Cys	Pro	Glu	Ala	Met	Asp	Leu	Gly	Thr
77			-	•	470	-	-			475					480
78	Leu	Ser	Glv	Ile	Gly	Thr	Leu	Asp	Gly	Phe	Arg	His	Arg	His	Pro
79					485			-		490	_				495
80	Asp	Glu	Ala	Ala	Phe	Phe	Asp	Thr	Ala	Ser	Thr	Gly	Lys	Thr	Phe
81					500		_			505					510
82	Pro	Glv	Phe	Phe	Ser	Pro	Met	Leu	Gly	Glu	Phe	Val	Ser	Glu	Thr
83					515				_	520					525
84	Glu	Ser	Arq	Gly	Ser	Glu	Ser	Gly	Ile	Phe	Thr	Asn	Thr	Lys	Glu
85			_	-	530			_		535					540
86	Ser	Ser	Ser	His	His	Pro	Gly	Ile	Ala	Glu	Phe	Pro	Ser	Arg	Gly
87					545		-			550					555
88	Lvs	Ser	Ser	Ser	Tyr	Ser	Lys	Gln	Phe	Thr	Ser	Ser	Thr	Ser	Tyr
89	-				560		_			565					570
90	Asn	Arq	Gly	Asp	Ser	Thr	Phe	Glu	Ser	Lys	Ser	Tyr	Lys	Met	Ala
91		,	-	-	575					580					585
92	Asp	Glu	Ala	Gly	Ser	Glu	Ala	Asp	His	Glu	Gly	Thr	His	Ser	Thr
93	-			-	590			-		595					600

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94 95	Lys Arg Gly His Ala Lys Ser Arg Pro Val Arg Gly Ile His Thr 605 610 615
96	Ser Pro Leu Gly Lys
97	620
99 <210>	SEQ ID NO: 2
100 <211>	LENGTH: 4
101 <212>	TYPE: PRT
102 <213>	ORGANISM: Artificial Sequence
103 <220>	
104 <223>	OTHER INFORMATION: Unknown. Obtained from a commercial source.
105 <400>	SEQUENCE: 2
106	Gly Pro Arg Pro